RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/583,840
Source:	IFWP
Date Processed by STIC:	07/06/2006

ENTERED



IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/583,840

DATE: 07/06/2006

TIME: 14:07:24

Input Set: A:\CMT-0072 sequence listing.txt
Output Set: N:\CRF4\07062006\J583840.raw

```
1 <110> APPLICANT: RHEE, Young Ha
              KIM, Young back
      2
              CHUNG, Chungwook
              YOON, Jooseog
              LG CHEM, LTD.
      7 <120> TITLE OF INVENTION: POLY(3-HYDROXYALKANOATE) BLOCK COPOLYMER HAVING SHAPE MEMORY
              EFFECT
     10 <130> FILE REFERENCE: CMT-0072
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/583,840
C--> 12 <141> CURRENT FILING DATE: 2006-06-22
     12 <150> PRIOR APPLICATION NUMBER: KR 10-2005-0059907
     13 <151> PRIOR FILING DATE: 2005-07-04
     15 <160> NUMBER OF SEQ ID NOS: 14
     17 <170> SOFTWARE: KopatentIn 1.71 and text editor
     19 <210> SEQ ID NO: 1
     20 <211> LENGTH: 18
     21 <212> TYPE: DNA
     22 <213> ORGANISM: Artificial Sequence
     24 <220> FEATURE:
     25 <223 > OTHER INFORMATION: Primer
     28 <400> SEQUENCE: 1
     29 ccgccstgsa tcaagtac
                                                                                    18
     32 <210> SEQ ID NO: 2
     33 <211> LENGTH: 20
     34 <212> TYPE: DNA
     35 <213> ORGANISM: Artificial Sequence
     37 <220> FEATURE:
     38 <223 > OTHER INFORMATION: Primer
     41 <400> SEQUENCE: 2
                                                                                    20
     42 gytsgtgsyg tcyycgttcc
     45 <210> SEQ ID NO: 3
     46 <211> LENGTH: 24
     47 <212> TYPE: DNA
     48 <213> ORGANISM: Artificial Sequence
     50 <220> FEATURE:
     51 <223> OTHER INFORMATION: Primer
     54 <400> SEQUENCE: 3
     55 caccatgctg agttgcgctc tagc
                                                                                    24
     58 <210> SEQ ID NO: 4
     59 <211> LENGTH: 27
     60 <212> TYPE: DNA
     61 <213> ORGANISM: Artificial Sequence
     63 <220> FEATURE:
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64 <223> OTHER INFORMATION: Primer	
67 <400> SEQUENCE: 4	2.7
68 tcadmsytty acrtarcgkc ctggygc	27
71 <210> SEQ ID NO: 5	
72 <211> LENGTH: 20	
73 <212> TYPE: DNA	
74 <213> ORGANISM: Artificial Sequence 76 <220> FEATURE:	
76 <2205 FEATURE. 77 <223> OTHER INFORMATION: Primer	
80 <400> SEQUENCE: 5	
81 gatcgatacc aatctcaccg	20
84 <210> SEQ ID NO: 6	
85 <211> LENGTH: 21	
86 <212> TYPE: DNA	
87 <213> ORGANISM: Artificial Sequence	
89 <220> FEATURE:	
90 <223> OTHER INFORMATION: Primer	
93 <400> SEQUENCE: 6	
94 caaagccagt ggttcgacgt a	21
97 <210> SEQ ID NO: 7	
98 <211> LENGTH: 19	
99 <212> TYPE: DNA	
100 <213> ORGANISM: Artificial Sequence	
102 <220> FEATURE:	
103 <223> OTHER INFORMATION: Primer	
106 <400> SEQUENCE: 7	
107 ctgctgaaac tgttggagc	19
110 <210> SEQ ID NO: 8	
111 <211> LENGTH: 47	
112 <212> TYPE: DNA	
113 <213> ORGANISM: Artificial Sequence	
115 <220> FEATURE:	
116 <223> OTHER INFORMATION: Primer	
119 <400> SEQUENCE: 8	
120 gggggtacca ataaggagat atacatatgg gtactgcgag caatgcg	47
123 <210> SEQ ID NO: 9	
124 <211> LENGTH: 28	
125 <212> TYPE: DNA	
126 <213 > ORGANISM: Artificial Sequence	
128 <220> FEATURE:	
129 <223> OTHER INFORMATION: Primer	
132 <400> SEQUENCE: 9	28
133 cccactagtt cagcgctcga tggccagc	20
136 <210> SEQ ID NO: 10	
137 <211> LENGTH: 28	
138 <212> TYPE: DNA	
139 <213> ORGANISM: Artificial Sequence	
141 <220> FEATURE:	
142 <223> OTHER INFORMATION: Primer	

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	.	too groupings to				
		<400> SEQUENCE: 10	0.0			
		gggcatatga cccagaagaa caacagcg	28			
		<210> SEQ ID NO: 11				
		<211> LENGTH: 39				
	_	<212> TYPE: DNA				
	152	2 <213> ORGANISM: Artificial Sequence				
	154	<220> FEATURE:				
	155	<223> OTHER INFORMATION: Primer				
	158	<400> SEQUENCE: 11				
		cccactagtt cadmscttya crtaacgtcc tggcgcygc	39			
	162	<210> SEQ ID NO: 12				
	163	<211> LENGTH: 756				
	_	<212> TYPE: DNA				
	165	<213> ORGANISM: Pseudomonas sp. HJ-2				
		<220> FEATURE:				
	168	<221> NAME/KEY: variation				
	169	<222> LOCATION: (482)				
	170	<223> OTHER INFORMATION: n=A, C, G or T				
		<400> SEQUENCE: 12				
		atgggtactg cgagcaatgc ggcacgtata gctctggtca ccggtggtat gggcggtatc	60			
	175	ggtacggcga tcagccagcg cctgcatcgg gatggcttca ccgtggtggt gggctgtaat	120			
	177	ccctactcca gccgcaaggc ttcctggatt gccacgcaac tcgaggcggg ctttcacttc	180			
	179	cactgcatcg actgcgacat caccgactgg gatagcaccc gccaggcctt cgacatggtg	240			
	181	cacgagactg tcggcccgat cgatgtattg gtcaacaatg ccggcatcac ccgcgacggc	300			
	183	actttccgca agatgtcccc ggaaaactgg aaggcggtga tcgataccaa tctcaccggc	360			
	185	ctgttcaaca caaccaagca ggtcatcgag ggcatgctgg ccaagggctg gggacgcgtc	420			
	187	atcaacatct cctcaatcaa tggccagcga ggccagttcg ggcagaccaa ctactccgcg	480			
W>	189	gncaaggctg gcattcatgg cttcagcatg gccttggccc gcgaggtgag tggcaagggc	540			
	191	gtgaccgtca atacggtttc ccctggctac atcaagaccg acatgaccgc ggcgattcgc	600			
		ccggacatcc tcgaagacat gattactggc attcccgtgg gccgtctcgg ccagcccgag	660			
	195	gagategeet egategtgge etggetggee teegateagt etgeetatge eaceggegee	720			
		gacttctcgg tgaatggcgg catgaacatg cagtga	756			
		<210> SEQ ID NO: 13				
		<211> LENGTH: 1179				
		<212> TYPE: DNA				
		<213> ORGANISM: Pseudomonas sp. HJ-2				
		<220> FEATURE:				
		<221> NAME/KEY: variation				
		<222> LOCATION: (207)				
		<223> OTHER INFORMATION: n=A, C, G or T				
	_	<220> FEATURE:				
		<221> NAME/KEY: variation				
		<222> LOCATION: (209)				
		<223> OTHER INFORMATION: n=A, C, G or T				
		<400> SEQUENCE: 13	~ ~			
		atgatcgaag tcgttatcgt cgccgccact cgcaccgcca tcggcgcttt ccaggggagc	60			
		ctggccggca ctcccgccgt tgaactgggc gccacggtga tccgccgcct gctcgaacag	120			
		accgctctgg atagcagtca ggtggatgaa gtgatactcg gccacgtact caccgccggt	180			
M>	223	gctggcagaa taccgctcgc caggcancng gtcatcgccg gcctgccaca cgccgtaccg	240			

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225	gcgatgaccc	tgaacaaggt	ctgtggctcc	ggcctgaaag	ccctgcacct	gggcgcccag	300
227	gccatccgct	gtggcgatgc	cgaggtggtg	attgccggtg	gcatggagaa	catgagcctg	360
229	tcgtcctatg	tcctgcccaa	ggcccgcacc	ggcctgcgca	tgggccacgc	gcagctggtc	420
231	gacagcatga	tcgtcgacgg	cctgtgggac	gccttcaacg	actaccacat	ggggatcact	480
233	gccgagaacc	tggtagacaa	gtacggcatc	agccgcgaag	cccaggacga	attcgccgcc	540
			ggccgccatc				600
237	ccggtgagca	ttccgcagcg	caagggcgag	gcgctgagct	tcgacaccga	cgaacagcca	660
		- -	gtcgctgggc				720
	· =		ttccagtctc				780
			gcttggtctg				840
			gatcatgggt				900
			tctggcagag				960
			gggtcaggag				1020
			cggccacccc				1080
			caggcgcgac		gcctcgctac	cctgtgtatc	1140
			gctggccatc	gagcgctga			1179
	<210> SEQ]						
	<211> LENGT						
	<212> TYPE:						
			omonas sp. I	HJ - 2			
	<400> SEQUE		.	+ ~ ~ + ~ ~ ~ ~ + ~	200000000	antagganga	60
			tgctcactac				120
	-		cttatacgtg			_	180
			acctgtcgag	_		_	240
			ccagcaattg	-			300
			ctggagcgaa				360
			gctgaaactg				420
			cttgatcgag				480
						cgccagccta	540
			tgccagtgac				600
						ggtactggaa	660
						ccagcggccg	720
						gcccgaaaac	780
						ctggcgcaac agtgatcagc	840
						gggtttctgc	900
	_					cgaccaggac	960
						gccgatcagc	1020
	_				-	ccatggtggc	1080
			tgaggacatg				1140
		•				gctggctctg	1200
						ttgctggtat	1260
						gtgcggcgtc	1320
						tgacgaccac	1380
			ctatgccagc				1440
						gaacaagcgc	1500
						gggagctcag	1560
						tgccggccca	1620
						acatgcgcca	1680
J I U	Jacangeerg		2222-426	22-2-2			

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320 ggacgttatg tgaagctatg a

1701

RAW SEQUENCE LISTING ERROR SUMMARY

PATENT APPLICATION: US/10/583,840

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Input Set: A:\CMT-0072 sequence listing.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:12; N Pos. 482 Seq#:13; N Pos. 207,209 VERIFICATION SUMMARY

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Input Set : A:\CMT-0072 sequence listing.txt
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L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:189 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:480 L:223 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:180